

10 Rec'd 30 JUL 2001

Attorney's Docket No.: 07898-070001 / PH-1046PCT-US

09/807123

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant : Yoshiharu Doi and Hiromi  
Matsusaki

Art Unit : Unknown  
Examiner : Unknown

Serial No. : 09/807,123

Filed : April 5, 2001

Title : METHOD OF PRODUCING COPOLYMER POLYESTER

Commissioner for Patents  
Washington, D.C. 20231

STATEMENT UNDER 37 CFR §1.821(f) AND (g)

I hereby state, as required by 37 C.F.R. §1.821(f), that the content of the paper and computer-readable copy of the Sequence Listing, submitted in accordance with 37 C.F.R. §§1.821(c) and (e), respectively, are the same.

I hereby state, as required by 37 C.F.R. §1.821(g), that the enclosed submission includes no new matter.

Respectfully submitted,

Date:

July 30, 2001

Mi K. Kim

Reg. No. 44,830

Fish & Richardson P.C.  
4350 La Jolla Village Drive, Suite 500  
San Diego, CA 92122  
Telephone: (858) 678-5070  
Facsimile: (858) 678-5099  
10125336.doc

CERTIFICATE OF MAILING BY EXPRESS MAIL

Express Mail Label No. EL558602025US

I hereby certify under 37 CFR §1.10 that this correspondence is being deposited with the United States Postal Service as Express Mail Post Office to Addressee with sufficient postage on the date indicated below and is addressed to the Commissioner for Patents, Washington, D.C. 20231.

July 30, 2001

Date of Deposit

Mike Augustine  
Signature

Mike Augustine

Typed or Printed Name of Person Signing Certificate



## UNITED STATES PATENT AND TRADEMARK OFFICE

Commissioner for Patents, Box PCT  
United States Patent and Trademark Office  
Washington, D.C. 20231  
www.uspto.gov

U.S. APPLICATION NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
09/807123	DOI	Y 07898-070001
INTERNATIONAL APPLICATION NO.		
PCT/JP00/05331		
I.A. FILING DATE		PRIORITY DATE
09 AUG 00		09 AUG 99

GREGORY P EINHORN  
FISH & RICHARDSON  
4350 LA JOLLA VILLAGE DRIVE SUITE 500  
SAN DIEGO, CA 92122

DATE MAILED: 2001

**NOTIFICATION TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS  
CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE  
DISCLOSURES**

Applicant has submitted papers under 35 U.S.C. 371 to enter the national stage in the United States of America. The items indicated below, however, are missing. The period within which to correct the deficiency noted below and avoid abandonment is set forth in the accompanying Notification.

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821-1.825 for the following reason(s):

- ☐ The application fails to comply with the requirements of 37 CFR 1.821-1.825.
- ☐ This application does not contain, a "Sequence Listing" as a separate part of the disclosure on paper copy or compact disc, as required by 37 CFR 1.821(c).
- ☐ A copy of the "Sequence Listing" in computer readable format has not been submitted as required by 37 CFR 1.821(e).
- ☒ A copy of the "Sequence Listing" in computer readable form has been submitted. The content of the computer readable form, however, does not comply with the requirements of 37 CFR 1.822 and/or 1.832, as indicated on the attached marked-up copy of the "Raw Sequence Listing."
- ☐ The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d).
- ☐ The paper copy or compact disc of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).
- ☐ Other: \_\_\_\_\_

**APPLICANT MUST PROVIDE:**

- ☒ An initial or substitute computer readable form (CRF) of the "Sequence Listing."
- ☐ An initial or substitute paper copy or compact disc of the "Sequence Listing," as well as an amendment directing its entry into the specification.
- ☒ A statement that the contents of the paper or compact disc and the computer readable form are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b) or 1.825(d).

**FOR QUESTIONS REGARDING COMPLIANCE WITH THESE REQUIREMENTS, PLEASE  
CALL:**

(703) 308-4216, for Rules interpretation,  
(703) 308-4212, for CRF submission help,  
(703) 287-0200, for PatentIn software help.

Fred Smith

Telephone: 703-305-3654

PCT09

## RAW SEQUENCE LISTING

DATE: 09/27/2001

PATENT APPLICATION: US/09/807,123

TIME: 12:00:14

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

2 <110> APPLICANT: Doi, Yoshiharu  
 3 Matsusaki, Hiromi  
 5 <120> TITLE OF INVENTION: METHOD OF PRODUCING COPOLYMER POLYESTER  
 8 <130> FILE REFERENCE: 07898-070001  
 10 <140> CURRENT APPLICATION NUMBER: 09/807,123  
 C--> 11 <141> CURRENT FILING DATE: 2001-07-31  
 13 <150> PRIOR APPLICATION NUMBER: PCT/JP00/05331  
 14 <151> PRIOR FILING DATE: 2000-08-09  
 16 <150> PRIOR APPLICATION NUMBER: JP 225102/1999  
 17 <151> PRIOR FILING DATE: 1999-08-09  
 19 <160> NUMBER OF SEQ ID NOS: 11  
 21 <170> SOFTWARE: PatentIn Ver. 2.0  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 1680  
 25 <212> TYPE: DNA  
 26 <213> ORGANISM: Pseudomonas sp. strain 61-3  
 28 <220> FEATURE:  
 29 <221> NAME/KEY: CDS  
 30 <222> LOCATION: (1)..(1677)  
 32 <400> SEQUENCE: 1

33	atg agt aac aag aat agc gat gac ttg aat cgt caa gcc tcg gaa aac	48
34	Met Ser Asn Lys Asn Ser Asp Asp Leu Asn Arg Gln Ala Ser Glu Asn	
35	1 5 10 15	
36	acc ttg ggg ctt aac cct gtc atc ggc ctg cgt gga aaa gat ctg ctg	96
37	Thr Leu Gly Leu Asn Pro Val Ile Gly Leu Arg Gly Lys Asp Leu Leu	
38	20 25 30	
39	act tct gcc cga atg gtt tta acc caa gcc atc aaa caa ccc att cac	144
40	Thr Ser Ala Arg Met Val Leu Thr Gln Ala Ile Lys Gln Pro Ile His	
41	35 40 45	
42	agc gtc aag cac gtc gcg cat ttt ggc atc gag ctg aag aac gtg atg	192
43	Ser Val Lys His Val Ala His Phe Gly Ile Glu Leu Lys Asn Val Met	
44	50 55 60	
45	ttt ggc aaa tcg aag ctg caa ccg gaa agc gat gac cgt cgt ttc aac	240
46	Phe Gly Lys Ser Lys Leu Gln Pro Glu Ser Asp Asp Arg Arg Phe Asn	
47	65 70 75 80	
48	gac ccc gcc tgg agt cag aac cca ctc tac aaa cgt tat cta caa acc	288
49	Asp Pro Ala Trp Ser Gln Asn Pro Leu Tyr Lys Arg Tyr Leu Gln Thr	
50	85 90 95	
51	tac ctg gcg tgg cgc aag gaa ctc cac gac tgg atc ggc aac agc aaa	336
52	Tyr Leu Ala Trp Arg Lys Glu Leu His Asp Trp Ile Gly Asn Ser Lys	
53	100 105 110	
54	ctg tcc gaa cag gac atc aat cgc gct cac ttc gtg atc acc ctg atg	384
55	Leu Ser Glu Gln Asp Ile Asn Arg Ala His Phe Val Ile Thr Leu Met	
56	115 120 125	
57	acc gaa gcc atg gcc ccg acc aac agt gcg gcc aat ccg gcg gcg gtc	432
58	Thr Glu Ala Met Ala Pro Thr Asn Ser Ala Ala Asn Pro Ala Ala Val	
59	130 135 140	

ENTERED

## RAW SEQUENCE LISTING

DATE: 09/27/2001

PATENT APPLICATION: US/09/807,123

TIME: 12:00:14

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

```

60 aaa cgc ttc ttc gaa acc ggc ggt aaa agc ctg ctc gac ggc ctc aca 480
61 Lys Arg Phe Phe Glu Thr Gly Gly Lys Ser Leu Leu Asp Gly Leu Thr
62 145 150 155 160
63 cat ctg gcc aag gac ctg gta aac aac ggc ggc atg ccg agc cag gtg 528
64 His Leu Ala Lys Asp Leu Val Asn Asn Gly Gly Met Pro Ser Gln Val
65 165 170 175
66 gac atg ggc gct ttc gaa gtc ggc aag agt ctg ggg acg act gaa ggt 576
67 Asp Met Gly Ala Phe Glu Val Gly Lys Ser Leu Gly Thr Thr Glu Gly
68 180 185 190
69 gca gtg gtt ttc cgc aac gac gtc ctc gaa ttg atc cag tac cgg ccg 624
70 Ala Val Val Phe Arg Asn Asp Val Leu Glu Leu Ile Gln Tyr Arg Pro
71 195 200 205
72 acc acc gaa cag gtg cat gag cga ccg ctg ctg gtg gtc cca ccg cag 672
73 Thr Thr Glu Gln Val His Glu Arg Pro Leu Leu Val Val Pro Pro Gln
74 210 215 220
75 atc aac aag ttt tat gtg ttt gac ctg agc ccg gat aaa agc ctg gcg 720
76 Ile Asn Lys Phe Tyr Val Phe Asp Leu Ser Pro Asp Lys Ser Leu Ala
77 225 230 235 240
78 cgc ttc tgc ctg agc aac aac cag caa acc ttt atc gtc agc tgg cgc 768
79 Arg Phe Cys Leu Ser Asn Asn Gln Gln Thr Phe Ile Val Ser Trp Arg
80 245 250 255
81 aac ccg acc aag gcc cag cgt gag tgg ggt ctg tcg act tac atc gat 816
82 Asn Pro Thr Lys Ala Gln Arg Glu Trp Gly Leu Ser Thr Tyr Ile Asp
83 260 265 270
84 gcg ctc aaa gaa gcc gtc gac gta gtt tcc gcc atc acc ggc agc aaa 864
85 Ala Leu Lys Glu Ala Val Asp Val Val Ser Ala Ile Thr Gly Ser Lys
86 275 280 285
87 gac atc aac atg ctc ggc gcc tgc tcc ggt ggc att acc tgc acc gcg 912
88 Asp Ile Asn Met Leu Gly Ala Cys Ser Gly Gly Ile Thr Cys Thr Ala
89 290 295 300
90 ctg ctg ggt cac tac gcc gct ctc ggc gag aag aag gtc aat gcc ctg 960
91 Leu Leu Gly His Tyr Ala Ala Leu Gly Glu Lys Lys Val Asn Ala Leu
92 305 310 315 320
93 acc ctt ttg gtc agc gtg ctc gac acc acc ctc gac tcc cag gtt gca 1008
94 Thr Leu Leu Val Ser Val Leu Asp Thr Thr Leu Asp Ser Gln Val Ala
95 325 330 335
96 ctg ttc gtc gat gag aaa acc ctg gaa gct gcc aag cgt cac tcg tat 1056
97 Leu Phe Val Asp Glu Lys Thr Leu Glu Ala Ala Lys Arg His Ser Tyr
98 340 345 350
99 cag gcc ggc gtg ctg gaa ggc cgc gac atg gcc aaa gtc ttc gcc tgg 1104
100 Gln Ala Gly Val Leu Glu Gly Arg Asp Met Ala Lys Val Phe Ala Trp
101 355 360 365
102 atg cgc cct aac gac ctg atc tgg aac tac tgg gtc aac aac tac ctg 1152
103 Met Arg Pro Asn Asp Leu Ile Trp Asn Tyr Trp Val Asn Asn Tyr Leu
104 370 375 380
105 ctg ggt aac gag cca ccg gtc ttc gac att ctt ttc tgg aac aac gac 1200
106 Leu Gly Asn Glu Pro Pro Val Phe Asp Ile Leu Phe Trp Asn Asn Asp
107 385 390 395 400
108 acc acc ccg ttg cct gct gcg ttc cac ggc gat ctg atc gaa atg ttc 1248

```

## RAW SEQUENCE LISTING

DATE: 09/27/2001

PATENT APPLICATION: US/09/807,123

TIME: 12:00:14

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

```

109 Thr Thr Arg Leu Pro Ala Ala Phe His Gly Asp Leu Ile Glu Met Phe
110          405          410          415
111 aaa aat aac cca ctg gtg cgc gcc aat gca ctc gaa gtg agc ggc acg 1296
112 Lys Asn Asn Pro Leu Val Arg Ala Asn Ala Leu Glu Val Ser Gly Thr
113          420          425          430
114 ccg atc gac ctc aaa cag gtc act gcc gac atc tac tcc ctg gcc ggc 1344
115 Pro Ile Asp Leu Lys Gln Val Thr Ala Asp Ile Tyr Ser Leu Ala Gly
116          435          440          445
117 acc aac gat cac atc acg ccc tgg aag tct tgc tac aag tcg gcg caa 1392
118 Thr Asn Asp His Ile Thr Pro Trp Lys Ser Cys Tyr Lys Ser Ala Gln
119          450          455          460
120 ctg ttc ggt ggc aag gtc gaa ttc gtg ctg tcc agc agt ggg cat atc 1440
121 Leu Phe Gly Gly Lys Val Glu Phe Val Leu Ser Ser Ser Gly His Ile
122 465          470          475          480
123 cag agc att ctg aac ccg ccg ggc aat ccg aaa tca cgt tac atg acc 1488
124 Gln Ser Ile Leu Asn Pro Pro Gly Asn Pro Lys Ser Arg Tyr Met Thr
125          485          490          495
126 agc acc gac atg cca gcc acc gcc aac gag tgg caa gaa aac tca acc 1536
127 Ser Thr Asp Met Pro Ala Thr Ala Asn Glu Trp Gln Glu Asn Ser Thr
128          500          505          510
129 aag cac acc gac tcc tgg tgg ctg cac tgg cag gcc tgg cag gcc gag 1584
130 Lys His Thr Asp Ser Trp Trp Leu His Trp Gln Ala Trp Gln Ala Glu
131          515          520          525
132 cgc tcg ggc aaa ctg aaa aag tcc ccg acc agc ctg ggc aac aag gcc 1632
133 Arg Ser Gly Lys Leu Lys Lys Ser Pro Thr Ser Leu Gly Asn Lys Ala
134          530          535          540
135 tat ccg tca gga gaa gcc gcg ccg ggc acg tat gtg cat gaa cgt taa 1680
136 Tyr Pro Ser Gly Glu Ala Ala Pro Gly Thr Tyr Val His Glu Arg
137 545          550          555
139 <210> SEQ ID NO: 2
140 <211> LENGTH: 559
141 <212> TYPE: PRT
142 <213> ORGANISM: Pseudomonas sp. strain 61-3
144 <400> SEQUENCE: 2
145 Met Ser Asn Lys Asn Ser Asp Asp Leu Asn Arg Gln Ala Ser Glu Asn
146 1          5          10          15
147 Thr Leu Gly Leu Asn Pro Val Ile Gly Leu Arg Gly Lys Asp Leu Leu
148          20          25          30
149 Thr Ser Ala Arg Met Val Leu Thr Gln Ala Ile Lys Gln Pro Ile His
150          35          40          45
151 Ser Val Lys His Val Ala His Phe Gly Ile Glu Leu Lys Asn Val Met
152          50          55          60
153 Phe Gly Lys Ser Lys Leu Gln Pro Glu Ser Asp Asp Arg Arg Phe Asn
154 65          70          75          80
155 Asp Pro Ala Trp Ser Gln Asn Pro Leu Tyr Lys Arg Tyr Leu Gln Thr
156          85          90          95
157 Tyr Leu Ala Trp Arg Lys Glu Leu His Asp Trp Ile Gly Asn Ser Lys
158          100          105          110
159 Leu Ser Glu Gln Asp Ile Asn Arg Ala His Phe Val Ile Thr Leu Met

```

## RAW SEQUENCE LISTING

DATE: 09/27/2001

PATENT APPLICATION: US/09/807,123

TIME: 12:00:14

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

```

160          115          120          125
161 Thr Glu Ala Met Ala Pro Thr Asn Ser Ala Ala Asn Pro Ala Ala Val
162          130          135          140
163 Lys Arg Phe Phe Glu Thr Gly Gly Lys Ser Leu Leu Asp Gly Leu Thr
164 145          150          155          160
165 His Leu Ala Lys Asp Leu Val Asn Asn Gly Gly Met Pro Ser Gln Val
166          165          170          175
167 Asp Met Gly Ala Phe Glu Val Gly Lys Ser Leu Gly Thr Thr Glu Gly
168          180          185          190
169 Ala Val Val Phe Arg Asn Asp Val Leu Glu Leu Ile Gln Tyr Arg Pro
170          195          200          205
171 Thr Thr Glu Gln Val His Glu Arg Pro Leu Leu Val Val Pro Pro Gln
172          210          215          220
173 Ile Asn Lys Phe Tyr Val Phe Asp Leu Ser Pro Asp Lys Ser Leu Ala
174 225          230          235          240
175 Arg Phe Cys Leu Ser Asn Asn Gln Gln Thr Phe Ile Val Ser Trp Arg
176          245          250          255
177 Asn Pro Thr Lys Ala Gln Arg Glu Trp Gly Leu Ser Thr Tyr Ile Asp
178          260          265          270
179 Ala Leu Lys Glu Ala Val Asp Val Val Ser Ala Ile Thr Gly Ser Lys
180          275          280          285
181 Asp Ile Asn Met Leu Gly Ala Cys Ser Gly Gly Ile Thr Cys Thr Ala
182          290          295          300
183 Leu Leu Gly His Tyr Ala Ala Leu Gly Glu Lys Lys Val Asn Ala Leu
184 305          310          315          320
185 Thr Leu Leu Val Ser Val Leu Asp Thr Thr Leu Asp Ser Gln Val Ala
186          325          330          335
187 Leu Phe Val Asp Glu Lys Thr Leu Glu Ala Ala Lys Arg His Ser Tyr
188          340          345          350
189 Gln Ala Gly Val Leu Glu Gly Arg Asp Met Ala Lys Val Phe Ala Trp
190          355          360          365
191 Met Arg Pro Asn Asp Leu Ile Trp Asn Tyr Trp Val Asn Asn Tyr Leu
192          370          375          380
193 Leu Gly Asn Glu Pro Pro Val Phe Asp Ile Leu Phe Trp Asn Asn Asp
194 385          390          395          400
195 Thr Thr Arg Leu Pro Ala Ala Phe His Gly Asp Leu Ile Glu Met Phe
196          405          410          415
197 Lys Asn Asn Pro Leu Val Arg Ala Asn Ala Leu Glu Val Ser Gly Thr
198          420          425          430
199 Pro Ile Asp Leu Lys Gln Val Thr Ala Asp Ile Tyr Ser Leu Ala Gly
200          435          440          445
201 Thr Asn Asp His Ile Thr Pro Trp Lys Ser Cys Tyr Lys Ser Ala Gln
202          450          455          460
203 Leu Phe Gly Gly Lys Val Glu Phe Val Leu Ser Ser Ser Gly His Ile
204 465          470          475          480
205 Gln Ser Ile Leu Asn Pro Pro Gly Asn Pro Lys Ser Arg Tyr Met Thr
206          485          490          495
207 Ser Thr Asp Met Pro Ala Thr Ala Asn Glu Trp Gln Glu Asn Ser Thr
208          500          505          510

```

## RAW SEQUENCE LISTING

DATE: 09/27/2001

PATENT APPLICATION: US/09/807,123

TIME: 12:00:14

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

```

209 Lys His Thr Asp Ser Trp Trp Leu His Trp Gln Ala Trp Gln Ala Glu
210          515          520          525
211 Arg Ser Gly Lys Leu Lys Lys Ser Pro Thr Ser Leu Gly Asn Lys Ala
212          530          535          540
213 Tyr Pro Ser Gly Glu Ala Ala Pro Gly Thr Tyr Val His Glu Arg
214 545          550          555
216 <210> SEQ ID NO: 3
217 <211> LENGTH: 1683
218 <212> TYPE: DNA
219 <213> ORGANISM: Pseudomonas sp. strain 61-3
W--> 220 <220> FEATURE:
221 <221> NAME/KEY: CDS
222 <222> LOCATION: (1)..(1680)
224 <400> SEQUENCE: 3
225 atg aga gag aaa cca acg ccg ggc ttg ctg ccc aca ccc gcg acg ttc 48
226 Met Arg Glu Lys Pro Thr Pro Gly Leu Leu Pro Thr Pro Ala Thr Phe
227 1          5          10          15
228 atc aac gct cag agt gcg att acc ggt ctg cgc ggc cgg gat ctg ttc 96
229 Ile Asn Ala Gln Ser Ala Ile Thr Gly Leu Arg Gly Arg Asp Leu Phe
230          20          25          30
231 tcg acc ctg cgc agc gtg gcc gcc cac ggc ctg cgt cac ccg gtg cgc 144
232 Ser Thr Leu Arg Ser Val Ala Ala His Gly Leu Arg His Pro Val Arg
233          35          40          45
234 agc gcc cgt cat gtt ctg gca ctg ggc ggc cag ttg ggc cgc gtg ctg 192
235 Ser Ala Arg His Val Leu Ala Leu Gly Gly Gln Leu Gly Arg Val Leu
236          50          55          60
237 ctg ggc gaa acg ctg cac acg ccg aac ccg aaa gac aat cgc ttt gcg 240
238 Leu Gly Glu Thr Leu His Thr Pro Asn Pro Lys Asp Asn Arg Phe Ala
239 65          70          75          80
240 gac ccg acc tgg aga ctg aat ccg ttt tac cgg cgc agc ctg cag gcc 288
241 Asp Pro Thr Trp Arg Leu Asn Pro Phe Tyr Arg Arg Ser Leu Gln Ala
242          85          90          95
243 tat ctg agc tgg cag aaa cag gtc aaa agc tgg atc gat gaa agc ggc 336
244 Tyr Leu Ser Trp Gln Lys Gln Val Lys Ser Trp Ile Asp Glu Ser Gly
245          100          105          110
246 atg agt gac gat gac cgc gcc cgc gcg cat ttc gtc ttc gca ctg ctc 384
247 Met Ser Asp Asp Asp Arg Ala Arg Ala His Phe Val Phe Ala Leu Leu
248          115          120          125
249 aat gac gcc gtg tcc ccc tcc aat acc ctg ctc aac ccg cta gcg atc 432
250 Asn Asp Ala Val Ser Pro Ser Asn Thr Leu Leu Asn Pro Leu Ala Ile
251          130          135          140
252 aag gag ctg ttc aac tcc ggt ggc aac agc ctg gtc cgc ggt ctc agc 480
253 Lys Glu Leu Phe Asn Ser Gly Gly Asn Ser Leu Val Arg Gly Leu Ser
254 145          150          155          160
255 cat tta ttc gac gac ctg atg cac aac aac ggg ctg ccc agt cag gtc 528
256 His Leu Phe Asp Asp Leu Met His Asn Asn Gly Leu Pro Ser Gln Val
257          165          170          175
258 acc aaa cac gcc ttc gag att ggc aag acc gtg gca acc acc gcc ggg 576
259 Thr Lys His Ala Phe Glu Ile Gly Lys Thr Val Ala Thr Thr Ala Gly

```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/807,123

DATE: 09/27/2001

TIME: 12:00:15

Input Set : A:\07898-070001doc.txt

Output Set: N:\CRF3\09272001\I807123.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:220 M:283 W: Missing Blank Line separator, <220> field identifier